

Database version 4.5
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Database has been searched using an index

Run on: September 4, 2002, 19:50:42 : Search time 13.139 seconds
(without alignments)
160,950 Million cell updates/sec

Index: Perfect score: 14
Sequence: 14

Scoring table:
Gapop 10.0 / Gapext 1.0

Searched: 1797656 seqs, 1045329254 residues

Total number of hits satisfying chosen parameters: 2545712

Minimum hit seq length: 0
Maximum hit seq length: 20000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listed first 45 summaries

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Database:
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2:  qb_dac:*
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31: qb_dac:*
32: qb_dac:*
33: qb_dac:*

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Prod. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No. Query Match Length Db. Hit Description

1	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
2	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
3	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
4	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
5	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
6	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
7	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
8	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
9	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
10	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
11	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
12	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
13	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
14	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
15	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
16	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
17	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
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19	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
20	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
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41	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
42	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
43	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
44	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999
45	42.3	15	6	AK03568	AK03568	Sequence 344 from Patent US 5809254.	15 bp	UNA	Linear	FAI 21-SEP-1999

ALIGNMENTS

RESULT 1	AK03568	15 bp	UNA	Linear	FAI 21-SEP-1999
DEFINITION	Sequence 344 from Patent US 5809254.				
ACCESSION	AK03568				
VERSION	AK03568.1				
KEYWORDS	GI:594173				
SOURCE	Unknown.				
ORGANISM	Unknown.				
REFERENCE	1 (Bases 1 to 15)				
AUTHORS	Draper, K.G.				
TITLE	Method and reagent for inhibiting hepatitis C virus replication				
JOURNAL	Patent: US 5809254-A 344 09 FEB 1999				
FEATURES	Location/Qualifiers				
SOURCE	1..15				
BASE COUNT	6 a 4 c 3 g 2 t				
ORIGIN	6 a 4 c 3 g 2 t				

Query Match 12.98% Score 62 DB 42 Length 15
Best Local Similarity 42.98% Prod. No. 100062

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ORIGIN
Query Match 42.9% Score 6: DB 6: Length 15:
Host Local Similarity 42.9% Prod. No. 1006:
Matches 6: Conservative 0: Mismatches 2: Indels 0: Gaps 0:
cy 1 ttatnnnnnnnn 14
    |||
    ||
db 14 ttgttgatgagct 1

RESULT 7
M005865/c 15 bp DNA linear PAT 31-JAN 2002
DEFINITION Brevi-factes (a) to detection of Mycobacteri
ACCESSION BD005865
VERSION BD005865.1 GI:18634236
KEYWORDS JP 2001501825-A/76.
SOURCE unidentified.
ORGANISM unidentified.
REFERENCE 1 (bases 1 to 15)
AUTHORS Stender,H., Lund,K. and Mollerup,T.A.
TITLE Novel probes for the detection of Mycobacteria
JOURNAL Patent: JP 2001501825-A 76 13-FEB-2001:
DAKO AS
COMMENT OS Unidentified
PN JP 2001501825-A/76
PD 13-FEB-2001
PE 03-OCT-1997 JP 1998517095
PR 04-OCT-1997 JP 1998517095
PS 05-MAV-1997 DK 0512/97
PI HENRIK STEINHE, KAREL LOMELTINA ANDRESEN BELMONT ET
CI 01/68,037814/00
CC Strandedness: Single;
CG Topology: Linear;
FH Key location/qualifiers
FI source 1..15
FEATURES
SOURCE location/qualifiers
    1..15 /organism="unidentified"
    /db_xref="taxon:32640"
BASE COUNT 6 a 7 c 1 g 1 t
ORIGIN
Query Match 42.9% Score 6: DB 6: Length 15:
Host Local Similarity 42.9% Prod. No. 1006:
Matches 6: Conservative 0: Mismatches 0: Indels 0: Gaps 0:
cy 1 ttatnnnnnnnn 14
    |||
    ||
db 14 ttgttgatgagct 1

RESULT 7
M005865/c 15 bp DNA linear PAT 31-JAN 2002
DEFINITION Brevi-factes (a) to detection of Mycobacteri
ACCESSION BD005865
VERSION BD005865.1 GI:18634236
KEYWORDS JP 2001501825-A/76.
SOURCE unidentified.
ORGANISM unidentified.
REFERENCE 1 (bases 1 to 15)
AUTHORS Stender,H., Lund,K. and Mollerup,T.A.
TITLE Novel probes for the detection of Mycobacteria
JOURNAL Patent: JP 2001501825-A 76 13-FEB-2001:
DAKO AS
COMMENT OS Unidentified
PN JP 2001501825-A/76
PD 13-FEB-2001
PE 03-OCT-1997 JP 1998517095
PR 04-OCT-1997 JP 1998517095
PS 05-MAV-1997 DK 0512/97
PI HENRIK STEINHE, KAREL LOMELTINA ANDRESEN BELMONT ET
CI 01/68,037814/00
CC Strandedness: Single;
CG Topology: Linear;
FH Key location/qualifiers
FI source 1..15
FEATURES
SOURCE location/qualifiers
    1..15 /organism="unidentified"
    /db_xref="taxon:32640"
BASE COUNT 6 a 7 c 1 g 1 t
ORIGIN
Query Match 42.9% Score 6: DB 6: Length 15:
Host Local Similarity 42.9% Prod. No. 1006:
Matches 6: Conservative 0: Mismatches 0: Indels 0: Gaps 0:
cy 1 ttatnnnnnnnn 14
    |||
    ||
db 14 ttgttgatgagct 1

```

```

unclassified.
REFERENCE 1 (bases 1 to 15)
AUTHORS Jerold,B.D.,Juli, and Harris,K.D.Y.Y.
TITLE Detection of Neisseria gonorrhoeae by amplifying and detecting
nucleic acid of Neisseria gonorrhoeae
JOURNAL Patent: JP 1999225781 A 11 24 AUG-1999:
DECTON DICKINSON & CO
OS Artificial Sequence
PN JP 1999225781-A/11
PD 24-AUG-1999
PE 30-OCT-1998 JP 1998309591
PR 04 NOV 1997 US 567363946
PI JEROLD B DAWWERTS,JAMES M HARRIS,KAREL DIRI VANSEN PO
CI 01/68,037814/00
CC Strandedness: Single;
CG Topology: Linear;
FH Key location/qualifiers
FI source 1..15
FEATURES
SOURCE location/qualifiers
    1..15 /organism="Artificial Sequence"
BASE COUNT 7 a 4 c 2 g 2 t
ORIGIN
Query Match 42.9% Score 6: DB 6: Length 15:
Host Local Similarity 42.9% Prod. No. 1006:
Matches 6: Conservative 0: Mismatches 0: Indels 0: Gaps 0:
cy 1 ttatnnnnnnnn 14
    |||
    ||
db 14 ttgttgatgagct 1

RESULT 9
E35697/c 15 bp DNA linear PAT 07-FEB-2001
DEFINITION Detection assay with the use of fluorescence and kit therefor.
ACCESSION E35697
VERSION E35697.1 GI:13019169
KEYWORDS JP 1999225799 A/13.
SOURCE unidentified.
ORGANISM unidentified.
REFERENCE 1 (bases 1 to 15)
AUTHORS Michael,C.L.G.G. and Yong.
TITLE Detection assay with the use of fluorescence and kit therefor
JOURNAL Patent: JP 1999225799 A 13 24-AUG-1999:
DECTON DICKINSON & CO
OS Artificial Sequence
PN JP 1999225799 A/13
PD 24-AUG-1999
PE 04-NOV-1998 JP 1998412790
PR 04 NOV 1997 US 567364020
PI MICHAEL C LITTLEBORN P YONG
CI 01/68,037814/00
CC Strandedness: Single;
CG Topology: Linear;
FH Key location/qualifiers
FI source 1..15
FEATURES
SOURCE location/qualifiers
    1..15 /organism="Artificial Sequence"
    /db_xref="taxon:32640"
BASE COUNT 7 a 4 c 2 g 2 t
ORIGIN
Query Match 42.9% Score 6: DB 6: Length 15:
Host Local Similarity 42.9% Prod. No. 1006:
Matches 6: Conservative 0: Mismatches 0: Indels 0: Gaps 0:
cy 1 ttatnnnnnnnn 14
    |||
    ||
db 14 ttgttgatgagct 1

```

```

db 14 TTTCATATTCAG 1
|||||
RESULT 10
LOCUS 157797
DEFINITION Sequences 444 from Patient US 5610954.
ACCESSION 5610954
VERSION 157797.1 GI:12482061
KEYWORDS
SOURCE
ORGANISM
REFERENCE
1 (bases 1 to 15)
AUTHORS
Draper, K.G.
TITLE
Enzymatic RNA molecule targeted against Hepatitis C virus
LOCATION/Qualifiers
1..15
FEATURES
BASE COUNT 6 a 4 c 3 g 2 t
ORIGIN
Query Match 42.9%; Score 6; DB 6; Length 15;
Best Local Similarity 42.9%; Pred. No. 10:06;
Matches 6; Conservative 0; Mismatches 0; Gaps 0;
db 15 TTTCATATTCAG 2
|||||
QUERY 1 tttgmmmmmmg 14
|||||
db 15 TTTCATATTCAG 2
|||||
RESULT 11
LOCUS AX132920
DEFINITION Sequence 4148 from Patent W0130362.
ACCESSION AX132920
VERSION AX132920.1 GI:14149240
KEYWORDS
SOURCE
ORGANISM
Homo sapiens
Eukaryotic Metazoa; Chordata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominoidea; Homo;
1 (bases 1 to 15)
AUTHORS
Kobayashi, M. and Iritani, R.
TITLE
Ribozyme therapy for the treatment of proliferative skin and eye
diseases
JOURNAL
Patient: W0130362 A 4148 03 MAY 2001;
IMMUSOL, INC. (US)
FEATURES
SOURCE
Location/Qualifiers
1..16
/organism "Homo sapiens"
/DB-access "W0130362"
/Note "Ribozyme fibrozyme recombinant site for cyclin B1"
BASE COUNT 0 a 4 3 c 7 g 6 t
ORIGIN
Query Match 42.9%; Score 6; DB 6; Length 16;
Best Local Similarity 42.9%; Pred. No. 10:06;
Matches 6; Conservative 0; Mismatches 0; Gaps 0;
db 6 TTTGCTGAGTGG 15
|||||
QUERY 1 tttgmmmmmmg 14
|||||
db 2 TTTCATATTCAG 15
|||||
RESULT 12
LOCUS A05414
DEFINITION Synthetic oligonucleotide primer.
ACCESSION A05414
VERSION A05414.1
AUTHORS
Patent: US 5617796 A 1212 06-oct 1998;
JOURNAL

```

```

ACCESSION A05414
VERSION A05414.1 GI:512617
KEYWORDS
SOURCE
ORGANISM
REFERENCE
1 (bases 1 to 17)
AUTHORS
Journal:
Patient: W0305144 A 28 17 MAY 1999;
Location/Qualifiers
1..17
/organism "Synthetic construct"
/DB-access "W0305144"
BASE COUNT 8 a 2 c 3 g 4 t
ORIGIN
Query Match 42.9%; Score 6; DB 6; Length 17;
Best Local Similarity 42.9%; Pred. No. 10:06;
Matches 6; Conservative 0; Mismatches 0; Gaps 0;
db 14 TTTCATATTCAG 1
|||||
QUERY 1 tttgmmmmmmg 14
|||||
db 14 TTTCATATTCAG 1
|||||
RESULT 13
LOCUS A09621
DEFINITION Oligonucleotide.
ACCESSION A09621
VERSION A09621.1 GI:490594
KEYWORDS
SOURCE
ORGANISM
Synthetic construct.
artificial sequence.
REFERENCE
1 (bases 1 to 17)
AUTHORS
Winter, G.P., Gressow, B. and Ward, E.S.
TITLE
Single domain ligands, receptors comprising said ligands, methods
for their production, and use of said ligands and receptors
JOURNAL
Patent: W0308684 A 48 16 MAY 1999;
MEDICAL RESEARCH COUNCIL.
FEATURES
SOURCE
Location/Qualifiers
1..17
/organism "Synthetic construct"
/DB-access "W0308684"
BASE COUNT 8 a 2 c 3 g 4 t
ORIGIN
Query Match 42.9%; Score 6; DB 6; Length 17;
Best Local Similarity 42.9%; Pred. No. 10:06;
Matches 6; Conservative 0; Mismatches 0; Gaps 0;
db 14 TTTCATATTCAG 1
|||||
QUERY 1 tttgmmmmmmg 14
|||||
db 14 TTTCATATTCAG 1
|||||
RESULT 14
LOCUS A046419
DEFINITION Sequence 1212 from Patient US 5817796.
ACCESSION A046419
VERSION A046419.1 GI:5967884
KEYWORDS
SOURCE
ORGANISM
Unknown.
Unclassified.
REFERENCE
1 (bases 1 to 17)
AUTHORS
Stichtomp, D.L., Draper, K.G., Mesquita, J. and Jarvis, J.
TITLE
Cyclic fibrozyne having 2'-5' linked adenylate residues
JOURNAL
Patent: US 5817796 A 1212 06-oct 1998;

```


CC The invention relates to the regulation of adenoviral packaging.
CC The method of the invention comprises propagating an adenoviral
CC vector containing a repressor binding site, in the absence of the
CC repressor. After propagation, vector packaging is repressed by the
CC appropriate repressor protein. The invention also encompasses an
CC adenoviral vector that includes an adenoviral packaging sequence
CC containing several copies of a chicken ovalbumin upstream promoter
CC transcription factor binding sites (AAV5997). Adenoviral vectors
CC containing repressor binding sites are used for DNA delivery, e.g., for
CC expression of a therapeutic protein, in genetic immunisation, or to
CC produce antiviral DNA or antisense RNA. Typical heterologous genes that
CC can be expressed include those for interferon-2, alpha-antitrypsin,
CC cyclic fibrosis transmembrane conductance regulator and coagulation
CC factor VIII. These vectors have very large capacity (up to 36 kb) for
CC foreign DNA and minimise the risk of generating replication competent
CC virus (since vector and helper virus can be designed such that they
CC have no overlapping packaging sequences that share primary control of
CC recombination). The presence of the repressor binding site allows
CC selective inhibition of vector packaging (10% packaging of one vector
CC in presence of another). Sequences AAV5996, AAV5997, AAV5998, AAV5999
CC represent a consensus of these A repeats.

SV Sequence 14 BP: 1 A; 3 C; 5 G; 5 T; 0 other;

Query Match 42.9%; Score 6; DB 21; Length 14;
Best Local Similarity 42.9%; Pred. No. 8, 1e+04;
Matches 6; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

QY 1 tttgmmmmmmnnnn 14
|||
||
Db 1 ttgtctagacccc 14

RESULT 4

AAV59895 standard; DNA: 14 BP.

AAV59895;

08-MAY-2000 (first entry)

Adenovirus minimal packaging element, A repeat, AV1.

Adenovirus: minimal packaging element, A repeat, sequence: binding site,
DNA delivery; ds.

Mastadenovirus.

W09951085-A2.

21-OCT-1999.

15-APR-1999; 99W0-US08294.

15-APR-1998; 98US-0081867.

05-JUN-1998; 98US-0088421.

(UNYNY) UNIV NEW YORK STATE RES FOUND.

Hearing P, Schmid SI, Ostapchuk PH, Er Turk E;

WPI: 2000-05267704

Regulating adenoviral packaging by incorporation of repressor binding
PT sites that allow selective suppression of packaging, used for gene
PI therapy

Disclosure: Page 15; 71PT; English.

The invention relates to the regulation of adenoviral packaging
CC The method of the invention comprises propagating an adenoviral

CC vector containing a repressor binding site, in the absence of the
CC repressor. After propagation, vector packaging is repressed by the
CC appropriate repressor protein. The invention also encompasses an
CC adenoviral vector that includes an adenoviral packaging sequence
CC containing several copies of a chicken ovalbumin upstream promoter
CC transcription factor binding sites (AAV5977). Adenoviral vectors
CC containing repressor binding sites are used for DNA delivery, e.g., for
CC expression of a therapeutic protein, in genetic immunisation, or to
CC produce antiviral DNA or antisense RNA. Typical heterologous genes that
CC can be expressed include those for interferon-2, alpha-antitrypsin,
CC cyclic fibrosis transmembrane conductance regulator and coagulation
CC factor VIII. These vectors have very large capacity (up to 36 kb) for
CC foreign DNA and minimise the risk of generating replication competent
CC virus (since vector and helper virus can be designed such that they
CC have no overlapping packaging sequences that share primary control of
CC recombination). The presence of the repressor binding site allows
CC selective inhibition of vector packaging (10% packaging of one vector
CC in presence of another). Sequences AAV5986, AAV5987, AAV5988, AAV5989
CC represent a consensus of these A repeats.

SV Sequence 14 BP: 2 A; 3 C; 5 G; 6 T; 0 other;

Query Match 42.9%; Score 6; DB 21; Length 14;
Best Local Similarity 42.9%; Pred. No. 8, 1e+04;
Matches 6; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

QY 1 tttgmmmmmmnnnn 14
|||
||
Db 1 ttgtctagacccc 14

RESULT 5

AAV6464270 standard; RNA: 15 BP.

AAV64642;

20-JUL-1999 (first entry)

Human R7-1 hammerhead ribozyme target SEQ ID NO:1274.

Arthritic condition; graft tolerance; immune response; target; cleavage;
KW hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; colibactinase;
KW streptococcus; streptococcus; streptococcus; streptococcus; streptococcus;
KW rheumatoid arthritis; autoimmune disease; allergy; inflammation;

diagnosis; ss.

Homo sapiens.

W09618736-A2.

20-JUN-1996.

22-NOV-1995; 95W0-0815516.

05-OCT-1995; 95US-0541165.

13-DEC-1994; 94US-0154620.

23-DEC-1994; 94US-0163253.

23-DEC-1994; 94US-0163254.

17-FEB-1995; 95US-0190850.

20-APR-1995; 95US-0126124.

02-MAY-1995; 95US-0123874.

04-MAY-1995; 95US-0134509.

07-JUL-1995; 95US-0000951.

07-JUL-1995; 95US-0000974.

07-AUG-1995; 95US-0512861.

(NAB) NABZTM; BACH INC.

Yoon E, Gershon E, Moshirpour F, Fink E, Gershon E;
PI Regulation of Fibroblast A, NABZ A, NABZ A, NABZ A

XX Sequence 15 bp: 5 A: 4 G: 4 C: 2 T: 0 other:

Query Match 42.9% Score 6: DB 21: Length 15:
 Best Local Similarity 42.9% Prod. No. 8.1e+04:
 Matches 6: Conservative 0: Mismatches 8: Indels 0: Gaps 0:

XX 1 TTTGAGAGAGAGAG 14
 DB 15 TTTGAGAGAGAG 2

RESULT 10

AAFG7497
 ID AAFG7497 standard: DNA, 15 bp.

XX AAFG7497

XX 11 JUN-2001 (first entry)

XX phage-opt vector amplifying primer.

XX (bacterial phage; pseudovirion; phagomid; pathogen; and bacterial).

XX (bacterial phage; pseudovirion; phagomid; pathogen; and bacterial).

XX Synthetic.

XX W (2001.21817.1).

XX 29 MAR-2001.

XX 22 SEP-2000: 2000W0-EP09.277.

XX 24 SEP-1999: 99EP-0402.448.

XX 03 NOV-1999: 99NS-0419.041.

XX (VIAA) VIAAMS INTERUNIVERSITAIR INST BIOTECHEMIE.

XX Myldegenius S. Silence K. Steyaert J. Torredo E.

XX W (2001.25795.256).

XX New genetically modified bacteriophage, pseudovirion or phagomid

XX capable of entering host cell by binding of its artificial ligand to

XX artificial receptor present on host cell, useful for eliminating

XX specific bacterial population -

XX Example 11: Page 45: 6pp: English.

XX The invention provides a genetically modified bacteriophage, pseudovirion

XX or phagomid (1) capable of entering a host cell by binding of its

XX artificial ligand (AL) to an artificial receptor (AR) present on the host

XX cell. (1) is useful for detecting and/or eliminating a specific bacterial

XX population, by AR-AL interaction, and to screen an antigen and/or

XX antibody library. (1) is useful for selecting AR-AL interactions. A kit

XX comprising (1) is useful for specific screening of a host cell, or

XX antibody treatment library or antibiotic sequences library. (1) is useful

XX for specific elimination of pathogenic bacteria, e.g., Aeromonas,

XX Enterococcus, Legionella, Listeria, Neisseria, etc and for screening a

XX host cell, displaying a bait against a library of bacteriophages/

XX pseudovirions/phagomids displaying the preys. The present sequence

XX represents a PCR primer for amplifying the phage-opt vector.

XX Sequence 15 bp: 5 A: 6 G: 2 C: 2 T: 0 other:

Query Match 42.9% Score 6: DB 22: Length 15:

Best Local Similarity 42.9% Prod. No. 8.1e+04:

Matches 6: Conservative 0: Mismatches 8: Indels 0: Gaps 0:

XX 1 TTTGAGAGAGAG 14

DB 15 TTTGAGAGAGAG 2

DB 14 TTTGAGAGAGAG 1

RESULT 11

AAFG4007
 ID AAFG4007 standard: DNA, 15 bp.

XX AAFG4007

XX 40-MAR-2001 (first entry)

XX IGFBP2 oligonucleotide #1147.

XX IGFBP2 oligonucleotide #1147.

XX Antisense therapy: antiproliferative and inflammatory; antiproliferative

XX cytoskeletal; dermatological; cardiac; vitreous; ophthalmic; keloid;

XX skin disorder; insulin-like growth factor 1 receptor; IGF 1; pituitary;

XX IGF binding protein; IGFBP-2; IGFBP-3; inflammation; psoriasis; pituitary;

XX growth factor mediated cell proliferation; ichthyosis; sorbitolase; tuba;

XX keratosis; neoplasia; scleroderma; warts; skin cancer; sclerotic disease;

XX hyperneovascular condition; hyperplasia; kidney disease;

XX neovascular condition of the retina; SS.

XX Homo sapiens.

XX W (2000.78341.1).

XX 21-JUN-2000: 2000W0-AD00693.

XX 21-JUN-1999: 99NS-0140.445.

XX (MORF) MORF-CH CHLORINE PEG INS.

XX W (2001.04142.705).

XX W (2001.04142.705).

XX Ameliorating the effects of a disorder, e.g., psoriasis, by

XX administering UV (ultraviolet) treatment (optional) and an antisense

XX nucleic acid that inhibits or reduces growth factor mediated cell

XX proliferation and/or inflammation -

XX Example 6: Page 41: 20pp: English.

XX The present invention relates to a method for ameliorating the effects

XX of skin disorders. The method comprises: (a) providing a skin with an

XX antisense oligonucleotide, (for insulin-like growth factor (IGF)-1

XX receptor, IGF binding protein (IGFBP-2 or IGFBP-3), which is capable of

XX inhibiting or reducing growth factor mediated cell proliferation,

XX inflammation and/or other disorders. The present sequence is an

XX oligonucleotide which can be used to design the antisense

XX oligonucleotides of the present invention (see AAFG4517) and

XX AAFG4517 #1513). The method is useful for ameliorating the effects of

XX psoriasis, ichthyosis, pityriasis, rube, folliculitis, seborrheic, keloids,

XX keratosis, neoplasia, scleroderma, warts, benign growths, cancers of the

XX retina, brain or skin, growth factor mediated malformations, other

XX sclerotic disease, kidney disease, hyperproliferation of the inside of

XX blood vessels or any other hyperplasia.

XX Sequence 15 bp: 4 A: 9 G: 1 C: 2 T: 0 other:

Query Match 42.9% Score 6: DB 22: Length 15:

Best Local Similarity 42.9% Prod. No. 8.1e+04:

Matches 6: Conservative 0: Mismatches 8: Indels 0: Gaps 0:

XX 1 TTTGAGAGAGAG 14

DB 15 TTTGAGAGAGAG 2

```

RESULT 12
AAFA47804/c
ID AAF46301 standard; DNA: 15 BP.
XX
AC AAF46301;
XX
DE 30-MAR-2001 (first entry)
XX
DE IGFBP2 oligonucleotide #1140.
XX
XX Antisense therapy: antiproliferative; antiinflammatory; antipsoriasis;
XX cytoskeletal; dermatological; cardiac; vitreous; ophthalmological; keloid;
XX skin disorder; insulin-like growth factor 1 receptor; psoriasis;
XX skin binding protein; IGFBP-2; IGFBP; inflammation; psoriasis; pilaris;
XX growth factor mediated cell proliferation; ichthyosis; scleroderma; ruba;
XX keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease;
XX hyperneovascular condition of the retina; kidney disease;
XX neovascular condition of the retina; ss.
XX
XX Homo sapiens.
XX
XX W0200078341-A1.
XX
XX 28-DEC-2000.
XX
XX 21-JUN-2000; 2000W0-A000693.
XX
XX 21-JUN-1999; 990S-0140345.
XX
XX (MORF) MORFCH CHILDENS RES INST.
XX
XX Wright CJ, Werther GA, Edmondson SK;
XX
XX WPI: 2001-041421/05.
XX
XX Ameliorating the effects of a disorder, e.g. psoriasis, by
XX administering UV (ultra-violet) treatment (optional) and an antisense
XX nucleic acid that inhibits or reduces growth factor mediated cell
XX proliferation and/or inflammation -
XX
XX Example 6; Page 41; 201pp; English.
XX
XX The present invention relates to a method for ameliorating the effects
XX of skin disorders. The method comprises contacting the skin with an
XX antisense oligonucleotide, (for insulin-like growth factor [IGF] 1
XX receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of
XX inhibiting or reducing growth factor mediated cell proliferation,
XX inflammation and/or other disorders. The present sequence is an
XX oligonucleotide which can be used to design the antisense
XX oligonucleotides of the present invention (see AAF45151 and
XX AAF45153-F45161). The method is useful for ameliorating the effects of
XX psoriasis, ichthyosis, pityriasis, ruba, pilaris, scleroderma, keloids,
XX keratosis, neoplasia, scleroderma, warts, benign growths, cancers of the
XX skin, a hyperneovascular condition such as a neovascular condition of the
XX retina, brain or skin, growth factor mediated malignancies, other
XX sclerotic disease, kidney disease, hyperproliferation of the inside of
XX blood vessels or any other hyperplasia.
XX
XX Sequence 15 BP; 3 A; 9 C; 1 G; 2 T; 0 other;
XX
XX
XX Query Match 42.9%; Score 6; DB 22; Length 15;
XX Best Local Similarity 42.9%; Pred. No. 8,1e+04;
XX Matches 6; Conservative 0; Mismatches 8; Indels 0; Gaps 0;
XX
XX 1 ttgmmmmmmmm 14
XX |||||
XX ||
XX 14 TTGGAGAGGCGG 1
XX
XX
XX RESULT 13
XX AAF47804/c
XX ID AAF47804 standard; DNA: 15 BP.
XX
XX
XX

```

```

XX
XX AAF47804;
XX
XX 30-MAR-2001 (first entry)
XX
XX IGFBP3 oligonucleotide #1224.
XX
XX
XX Antisense therapy: antiproliferative; antiinflammatory; antipsoriasis;
XX cytoskeletal; dermatological; cardiac; vitreous; ophthalmological; keloid;
XX skin disorder; insulin-like growth factor 1 receptor; IGFBP-1; pityriasis;
XX IGF binding protein; IGFBP-2; IGFBP; inflammation; psoriasis; pilaris;
XX growth factor mediated cell proliferation; ichthyosis; scleroderma; ruba;
XX keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease;
XX hyperneovascular condition of the retina; kidney disease;
XX neovascular condition of the retina; ss.
XX
XX Homo sapiens.
XX
XX W0200078341-A1.
XX
XX 28-DEC-2000.
XX
XX 21-JUN-2000; 2000W0-A000693.
XX
XX 21-JUN-1999; 990S-0140345.
XX
XX (MORF) MORFCH CHILDENS RES INST.
XX
XX Wright CJ, Werther GA, Edmondson SK;
XX
XX WPI: 2001-041421/05.
XX
XX Ameliorating the effects of a disorder, e.g. psoriasis, by
XX administering UV (ultra-violet) treatment (optional) and an antisense
XX nucleic acid that inhibits or reduces growth factor mediated cell
XX proliferation and/or inflammation -
XX
XX Example 7; Page 52; 201pp; English.
XX
XX The present invention relates to a method for ameliorating the effects
XX of skin disorders. The method comprises contacting the skin with an
XX antisense oligonucleotide, (for insulin-like growth factor [IGF]-1
XX receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of
XX inhibiting or reducing growth factor mediated cell proliferation,
XX inflammation and/or other disorders. The present sequence is an
XX oligonucleotide which can be used to design the antisense
XX oligonucleotides of the present invention (see AAF45151 and
XX AAF45153-F45161). The method is useful for ameliorating the effects of
XX psoriasis, ichthyosis, pityriasis, ruba, pilaris, scleroderma, keloids,
XX keratosis, neoplasia, scleroderma, warts, benign growths, cancers of the
XX skin, a hyperneovascular condition such as a neovascular condition of the
XX retina, brain or skin, growth factor mediated malignancies, other
XX sclerotic disease, kidney disease, hyperproliferation of the inside of
XX blood vessels or any other hyperplasia.
XX
XX Sequence 15 BP; 4 A; 5 C; 1 G; 5 T; 0 other;
XX
XX
XX Query Match 42.9%; Score 6; DB 22; Length 15;
XX Best Local Similarity 42.9%; Pred. No. 8,1e+04;
XX Matches 6; Conservative 0; Mismatches 8; Indels 0; Gaps 0;
XX
XX 1 ttgmmmmmmmm 14
XX |||||
XX ||
XX 15 TTGGAGAGGCGG 2
XX
XX
XX RESULT 14
XX AAF47805/c
XX ID AAF47805 standard; DNA: 15 BP.
XX
XX
XX AAF47805;
XX
XX

```


GenCore version 4.5
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nm nucleic nucleic search, using SW model

Run on: September 22, 2002, 22:51:17 : Search time 42.9 seconds
(without alignments)
80,160 Million cell updates/sec

Title: US-09-530-935 1

Sequence: 14 Nucleotide length 14

Scoring table: IDENTITY_NUC

Gapop 10.0 / Gapext 1.0

Searches: 38353 seqs, 12816752 residues

Total number of hits satisfying chosen parameters: 767056

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database: Issued_Patents_NA:
1: /path/to/database/2/Issued_Patents_NA.seq
2: /path/to/database/2/Issued_Patents_NA.seq
3: /path/to/database/2/Issued_Patents_NA.seq
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SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	6	42.9	15	US-08-182-968A-334	Sequence 334, App
2	6	42.9	15	US-08-774-206A-334	Sequence 334, App
3	6	42.9	15	US-08-663-946-11	Sequence 11, Appl
4	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
5	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
6	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
7	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
8	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
9	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
10	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
11	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
12	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
13	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
14	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
15	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
16	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
17	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
18	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
19	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
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21	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
22	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
23	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
24	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
25	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
26	6	42.9	15	US-08-963-946-11	Sequence 11, Appl
27	6	42.9	15	US-08-963-946-11	Sequence 11, Appl

ALIGNMENTS

28	6	42.9	18	US-09-194-277A-42	Sequence 42, Appl
29	6	42.9	18	PCT-US95-13142-30	Sequence 40, Appl
30	6	42.9	19	US-08-093-144-14	Sequence 14, Appl
31	6	42.9	19	US-09-031-442A-12	Sequence 12, Appl
32	6	42.9	19	US-08-599-507-4	Sequence 4, Appl
33	6	42.9	19	US-08-081-576-6	Sequence 6, Appl
34	6	42.9	19	US-09-384-305-12	Sequence 12, Appl
35	6	42.9	19	US-09-258-377-12	Sequence 12, Appl
36	6	42.9	19	US-08-274-606-5	Sequence 5, Appl
37	6	42.9	19	US-08-771-190-5	Sequence 5, Appl
38	6	42.9	19	US-09-189-294B-1	Sequence 1, Appl
39	6	42.9	19	US-09-189-294B-3	Sequence 3, Appl
40	6	42.9	19	US-09-189-294B-4	Sequence 4, Appl
41	6	42.9	19	PCT-US94-07955-5	Sequence 5, Appl
42	6	42.9	19	US-09-690-18	Sequence 18, Appl
43	6	42.9	20	US-07-621-001A-10	Sequence 10, Appl
44	6	42.9	20	US-08-096-182A-14	Sequence 14, Appl
45	6	42.9	20	US-08-474-542A-94	Sequence 94, Appl

RESULT 1
US-08-182-968A-334/c
Sequence 334, Application US/08182968A
Patent No. 5610054

GENERAL INFORMATION:
APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: IDENTIFYING HEMATITIS C
TYPE OF INVENTION: VITRO DETECTION
NUMBER OF SHEETS: 497
CORRESPONDENT ADDRESS:
ADDRESS: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
ATTORNEY, COUNSEL, OR AGENT: 08-093-144
FILING DATE: 13 JANUARY 1994
PRIORITY APPLICATION DATA:
APPLICATION NUMBER: 07-092,888
FILING DATE: 14 MAY 1992
ATTORNEY, COUNSEL, OR AGENT: 08-093-144
NAME: Wadburg, Richard J.
RESIDENTIAL ADDRESS:
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STATE: California
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ZIP: 90062-1000
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
FAX: 67-3510
INFORMATION FOR SEQ ID NO: 434:
SEQUENCE CHARACTERISTICS:
LENGTH: 15
TYPE: nucleic acid
STRATEGY: single
TOPOLOGY: linear
US-08-182-968A-334

Query Match: 42.9% Score: 63 DB ID: Length: 15
Best Local Similarity: 42.9% Pred. No. 1004
Ref. No. 1004 Pred. No. 1004
Ref. No. 1004 Pred. No. 1004

14 01:11:47:AR010001

Search completed: September 4, 2002, 2:43:59:29
Job Number: 8642, Size:
